

**Amendment to the claims**

1. (Currently amended) A method of tag-directed synthesis of a plurality of compounds, comprising:

(a) forming a first group of subsets of nucleic acid tags ~~for participating in a first synthetic reaction step,~~ where the nucleic acid tags in each subset each has a selected one of a plurality of different first hybridization sequences, a mixture of different second hybridization sequences, and a chemical reaction site, ~~by contacting said nucleic acid tags with a plurality of first immobilized nucleotide sequences, each designed to capture a subset of said tags by hybridization between one of said tag first hybridization sequences and the associated first immobilized sequence;~~

(b) ~~carrying out the first synthetic step by~~ reacting the chemical reaction sites in each of the subsets formed in (a) with a selected ~~one of a plurality of first reagents~~ first reagent, thereby to ~~convert the reaction site in each tag to form~~ a reagent-specific compound intermediate on the associated ~~sequence-tag~~ in each subset[.];

(c) forming a second group of subsets of the reacted nucleic acid tags ~~formed in step (b), for participation in a second synthetic reaction step, where the tags in each subset each have a selected one of a plurality of different second hybridization sequences, and a mixture of different first hybridization sequences~~ ~~by contacting said reacted nucleic acid tags with a plurality of second immobilized nucleotide sequences, each designed to capture a subset of said tags by hybridization between one of said tag second hybridization sequences and the associated second immobilized sequence;~~ and

(d) ~~carrying out the second synthetic step by~~ reacting the compound intermediates in the ~~associated tags~~ sequences in each of the subsets formed in (c) with a selected ~~one of a plurality of second reagents~~ first reagent, ~~whereby the nucleic acid tags direct the synthesis of the compounds.~~

2. (Cancelled)

3. (Currently amended) The method of claim 1, for use in forming a plurality of oligomers with different subunit sequences, wherein ~~the plurality of first and second reagents in steps (b) and (d) include different oligomer subunits each of said reacting steps includes adding a selected subunit to each of the subsets of nucleic acid tags under conditions effective to add that subunit to the chemical reaction site or last-added subunit carried thereon.~~

4. (Currently Amended) The method of claim 1, ~~for use in forming a plurality of compounds with different substituents, wherein the plurality of first and second reagents in steps (b) and (d) include different compound substituents each of said reacting steps includes adding a selected chemical substituent to each of the subsets of nucleic acid tags under conditions effective to add that substituent to the chemical reaction site or last-added substituent carried thereon.~~

5. (Currently Amended) The method of claim 1 for making a plurality of compounds requiring more than 2 synthetic steps wherein the nucleic acid tags include an additional step-specific subset of hybridization sequences for each synthetic step N greater than 2 and which further comprises, for each additional synthetic step N;

(e) forming an Nth group groups of subsets of reacted nucleic acid tags ~~for participating in the Nth reaction step, wherein the sequences in each subset each have a selected one of a plurality of hybridization sequences in a step-specific subset of nucleic acid tags, and a mixture of different other hybridization sequences~~ by contacting said nucleic acid tags with a plurality of Nth immobilized nucleotide sequences, each designed to capture a subset of said tags by hybridization between one of said tag Nth hybridization sequences and the associated Nth immobilized sequence;

(f) reacting the compound intermediates in the ~~associated tags~~ sequences in each of the subsets formed in (e) with a one of a plurality of Nth-reaction reagents ~~selected reagent;~~ and

(g) repeating steps (e) and (f) if necessary, until synthesis of the compounds is complete.

6. (Currently Amended) The method of claim 4 or 5 wherein each subset of nucleic acid tags includes at least 5 separate hybridization sequences.

7. (Original) The method of claim 1, wherein said nucleic acid tags within each subset further comprises for each subset of hybridization sequences, an adjacent spacer sequence separating that hybridization sequence from an adjacent one, each of said spacer sequences being the same for all subsets of nucleic acid tags and each hybridization sequence being different for each group of subsets of nucleic acid tags.

8. (Original) The method according to claim 1, for use in enriching the plurality of compounds in those having a desired compound activity, further comprising identifying from said plurality of compounds, one or more compounds having a desired activity to yield a subpopulation of nucleic acid tags, and using the subpopulation to carry out the method of claim 1.

9. (Currently Amended) The method according to claim 8, wherein said using includes;

amplifying said subpopulation of nucleic acid tags by ~~non-specific~~ polymerase chain reaction (PCR),  
adding a chemical reaction site, and  
using said amplified subpopulation having chemical reaction sites to carry out the method of claim 1.

10. (Original) The method according to claim 9, for use in producing new permutations of active compounds wherein said nucleic acid tags have one of a plurality of spacer sequences, each of said spacer sequences having a unique restriction enzyme site;

(e) identifying from said plurality of compounds, one or more compounds having a desired activity to yield a subpopulation of nucleic acid tags;

(f) treating said subpopulation of nucleic acid tags with one or more restriction enzymes under conditions effective to produce a partial digest;

(g) rejoining said partially digested nucleic acid tags;

(h) adding a new chemical reaction site to said partially digested nucleic acid tags and using the subpopulation to carry out the method of claim 1.

11. (Withdrawn) A library of nucleic acid tags for use in directing the synthesis of a plurality of compounds comprising a first segment having a selected one of a plurality of different first hybridization sequences, a second segment having a selected one of a plurality of different second hybridization sequences and a chemical reaction site wherein said first and second hybridization sequences are non-hybridizing with one another.

12. (Withdrawn) The library of nucleic acid tags according to claim 11, wherein said hybridization sequences are at least 10 nucleotides long.

13. (Withdrawn) The library of nucleic acid tags according to claim 11, for making a plurality of compounds requiring more than 2 synthetic steps wherein the nucleic acid sequences include an additional step-specific segment of hybridization sequences for each synthetic step greater than 2, wherein each sequence has a selected one of a plurality of hybridization sequences and said hybridization sequences are non-hybridizing with one another.

14. (Withdrawn) A combinatorial library of compounds, each combinatorial compound comprising a nucleic acid tag having a first segment with a selected one of a plurality of different first hybridization sequences, a second segment having a selected one of a plurality of different second hybridization sequences and a chemical reaction site wherein said first and second hybridization sequences are non-hybridizing with one another, and a compound formed on said chemical reaction site by step-wise chemical reaction which is directed by the associated nucleic acid tag.

15. (New) The method of claim 1, wherein each of said nucleotide sequences are each bound to the surface of a solid phase reagent.

16. (New) The method of claim 1, wherein said reacting steps (b) and (d) include first transferring the separate subsets of said tags from said immobilized sequences to solid support.